

SEQUENCE LISTING

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<120> EPITOPE SEQUENCES

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<151> 2002-09-06

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Pro Ala Asn Glu Tyr Ala Tyr Arg Arg Gly Ile Ala Glu Ala Val Gly		270
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	290	295
Leu Leu Glu Lys Met Gly Gly Ser Ala Pro Pro Asp Ser Ser Trp Arg		300
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Gly Ser Leu Lys Val Pro Tyr Asn Val Gly Pro Gly Phe Thr Gly Asn		320
	325	330
Phe Ser Thr Gln Lys Val Lys Met His Ile His Ser Thr Asn Glu Val		335
	340	345
Thr Arg Ile Tyr Asn Val Ile Gly Thr Leu Arg Gly Ala Val Glu Pro		350
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Asp Arg Tyr Val Ile Leu Gly Gly His Arg Asp Ser Trp Val Phe Gly		365
	370	375
Gly Ile Asp Pro Gln Ser Gly Ala Ala Val Val His Glu Ile Val Arg		380
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Ser Phe Gly Thr Leu Lys Lys Glu Gly Trp Arg Pro Arg Arg Thr Ile		400
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Glu Trp Ala Glu Glu Asn Ser Arg Leu Leu Gln Glu Arg Gly Val Ala		430
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Tyr Ile Asn Ala Asp Ser Ser Ile Glu Gly Asn Tyr Thr Leu Arg Val		445
	450	455
Asp Cys Thr Pro Leu Met Tyr Ser Leu Val His Asn Leu Thr Lys Glu		460
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Trp Thr Lys Lys Ser Pro Ser Pro Glu Phe Ser Gly Met Pro Arg Ile		495
	500	505
Ser Lys Leu Gly Ser Gly Asn Asp Phe Glu Val Phe Phe Gln Arg Leu		510
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Gly Ile Ala Ser Gly Arg Ala Arg Tyr Thr Lys Asn Trp Glu Thr Asn		525
	530	535
Lys Phe Ser Gly Tyr Pro Leu Tyr His Ser Val Tyr Glu Thr Tyr Glu		540
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Ala Gln Val Arg Gly Gly Met Val Phe Glu Leu Ala Asn Ser Ile Val		575
	580	585
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Met Lys Ala Ser Glu Lys Ile Phe Tyr
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Lys Met Lys Ala Ser Glu Lys Ile Phe Tyr
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Lys Ala Ser Glu Lys Ile Phe Tyr Val
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Met Lys Ala Ser Glu Lys Ile Phe Tyr Val
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Met Pro Glu Gly Asp Leu Val Tyr
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<210> 37
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<400> 37
Glu Gly Asp Leu Val Tyr Val Asn Tyr
1 5

<210> 38
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<400> 38
Pro Glu Gly Asp Leu Val Tyr Val Asn Tyr
1 5 10

<210> 39
<211> 10
<212> PRT
<213> Homo sapiens

<400> 39
Leu Val Tyr Val Asn Tyr Ala Arg Thr Glu
1 5 10

<210> 40
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<212> PRT
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<400> 40
 Val Asn Tyr Ala Arg Thr Glu Asp Phe
 1 5

<210> 41
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<400> 41
 Tyr Val Asn Tyr Ala Arg Thr Glu Asp Phe
 1 5 10

<210> 42
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<400> 42
 Asn Tyr Ala Arg Thr Glu Asp Phe Phe
 1 5

<210> 43
 <211> 8
 <212> PRT
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<400> 43
 Tyr Ala Arg Thr Glu Asp Phe Phe
 1 5

<210> 44
 <211> 9
 <212> PRT
 <213> Homo sapiens

<400> 44
 Arg Thr Glu Asp Phe Phe Lys Leu Glu
 1 5

<210> 45
 <211> 30
 <212> PRT
 <213> Homo sapiens

<400> 45
 Arg Gly Ile Ala Glu Ala Val Gly Leu Pro Ser Ile Pro Val His Pro
 1 5 10 15
 Ile Gly Tyr Tyr Asp Ala Gln Lys Leu Leu Glu Lys Met Gly
 20 25 30

<210> 46
<211> 25
<212> PRT
<213> Homo sapiens

<400> 46
Ile Ala Glu Ala Val Gly Leu Pro Ser Ile Pro Val His Pro Ile Gly
1 5 10 15
Tyr Tyr Asp Ala Gln Lys Leu Leu Glu
20 25

<210> 47
<211> 9
<212> PRT
<213> Homo sapiens

<400> 47
Leu Pro Ser Ile Pro Val His Pro Ile
1 5

<210> 48
<211> 10
<212> PRT
<213> Homo sapiens

<400> 48
Gly Leu Pro Ser Ile Pro Val His Pro Ile
1 5 10

<210> 49
<211> 9
<212> PRT
<213> Homo sapiens

<400> 49
Ile Gly Tyr Tyr Asp Ala Gln Lys Leu
1 5

<210> 50
<211> 10
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<400> 50
Pro Ile Gly Tyr Tyr Asp Ala Gln Lys Leu
1 5 10

<210> 51
<211> 9
<212> PRT
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<400> 51
Ser Ile Pro Val His Pro Ile Gly Tyr
1 5

<210> 52
<211> 10
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<213> Homo sapiens

<400> 52
Pro Ser Ile Pro Val His Pro Ile Gly Tyr
1 5 10

<210> 53
<211> 8
<212> PRT
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<400> 53
Ile Pro Val His Pro Ile Gly Tyr
1 5

<210> 54
<211> 9
<212> PRT
<213> Homo sapiens

<400> 54
Tyr Tyr Asp Ala Gln Lys Leu Leu Glu
1 5

<210> 55
<211> 27
<212> PRT
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<400> 55
Ser Ser Ile Glu Gly Asn Tyr Thr Leu Arg Val Asp Cys Thr Pro Leu
1 5 10 15
Met Tyr Ser Leu Val His Leu Thr Lys Glu Leu
20 25

<210> 56
<211> 9
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<400> 56
Ile Glu Gly Asn Tyr Thr Leu Arg Val
1 5

<210> 57
<211> 10
<212> PRT
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<400> 57
Ser Ile Glu Gly Asn Tyr Thr Leu Arg Val
1 5 10

<210> 58
<211> 8
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<400> 58
Glu Gly Asn Tyr Thr Leu Arg Val
1 5

<210> 59
<211> 9
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<400> 59
Thr Leu Arg Val Asp Cys Thr Pro Leu
1 5

<210> 60
<211> 10
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<400> 60
Tyr Thr Leu Arg Val Asp Cys Thr Pro Leu
1 5 10

<210> 61
<211> 9
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<400> 61
Leu Arg Val Asp Cys Thr Pro Leu Met
1 5

<210> 62
<211> 9
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<400> 62

Arg Val Asp Cys Thr Pro Leu Met Tyr
1 5

<210> 63
<211> 10
<212> PRT
<213> Homo sapiens

<400> 63
Leu Arg Val Asp Cys Thr Pro Leu Met Tyr
1 5 10

<210> 64
<211> 35
<212> PRT
<213> Homo sapiens

<400> 64
Phe Asp Lys Ser Asn Pro Ile Val Leu Arg Met Met Asn Asp Gln Leu
1 5 10 15
Met Phe Leu Glu Arg Ala Phe Ile Asp Pro Leu Gly Leu Pro Asp Arg
20 25 30
Pro Phe Tyr
35

<210> 65
<211> 22
<212> PRT
<213> Homo sapiens

<400> 65
Val Leu Arg Met Met Asn Asp Gln Leu Met Phe Leu Glu Arg Ala Phe
1 5 10 15
Ile Asp Pro Leu Gly Leu
20

<210> 66
<211> 9
<212> PRT
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<400> 66
Met Met Asn Asp Gln Leu Met Phe Leu
1 5

<210> 67
<211> 10
<212> PRT
<213> Homo sapiens

<400> 67
Arg Met Met Asn Asp Gln Leu Met Phe Leu

1 5 10

<210> 68
 <211> 9
 <212> PRT
 <213> Homo sapiens

<400> 68
 Arg Met Met Asn Asp Gln Leu Met Phe
 1 5

<210> 69
 <211> 17
 <212> PRT
 <213> Homo sapiens

<400> 69
 Met Leu Leu Ala Val Leu Tyr Cys Leu Leu Trp Ser Phe Gln Thr Ser
 1 5 10 15
 Ala

<210> 70
 <211> 661
 <212> PRT
 <213> Homo sapiens

<400> 70
 Met Asp Leu Val Leu Lys Arg Cys Leu Leu His Leu Ala Val Ile Gly
 1 5 10 15
 Ala Leu Leu Ala Val Gly Ala Thr Lys Val Pro Arg Asn Gln Asp Trp
 20 25 30
 Leu Gly Val Ser Arg Gln Leu Arg Thr Lys Ala Trp Asn Arg Gln Leu
 35 40 45
 Tyr Pro Glu Trp Thr Glu Ala Gln Arg Leu Asp Cys Trp Arg Gly Gly
 50 55 60
 Gln Val Ser Leu Lys Val Ser Asn Asp Gly Pro Thr Leu Ile Gly Ala
 65 70 75 80
 Asn Ala Ser Phe Ser Ile Ala Leu Asn Phe Pro Gly Ser Gln Lys Val
 85 90 95
 Leu Pro Asp Gly Gln Val Ile Trp Val Asn Asn Thr Ile Ile Asn Gly
 100 105 110
 Ser Gln Val Trp Gly Gly Gln Pro Val Tyr Pro Gln Glu Thr Asp Asp
 115 120 125
 Ala Cys Ile Phe Pro Asp Gly Gly Pro Cys Pro Ser Gly Ser Trp Ser
 130 135 140
 Gln Lys Arg Ser Phe Val Tyr Val Trp Lys Thr Trp Gly Gln Tyr Trp
 145 150 155 160
 Gln Val Leu Gly Gly Pro Val Ser Gly Leu Ser Ile Gly Thr Gly Arg
 165 170 175
 Ala Met Leu Gly Thr His Thr Met Glu Val Thr Val Tyr His Arg Arg
 180 185 190
 Gly Ser Arg Ser Tyr Val Pro Leu Ala His Ser Ser Ser Ala Phe Thr
 195 200 205

Ile	Thr	Asp	Gln	Val	Pro	Phe	Ser	Val	Ser	Val	Ser	Gln	Leu	Arg	Ala	210	215	220
Leu	Asp	Gly	Gly	Asn	Lys	His	Phe	Leu	Arg	Asn	Gln	Pro	Leu	Thr	Phe	225	230	235
Ala	Leu	Gln	Leu	His	Asp	Pro	Ser	Gly	Tyr	Leu	Ala	Glu	Ala	Asp	Leu	245	250	255
Ser	Tyr	Thr	Trp	Asp	Phe	Gly	Asp	Ser	Ser	Gly	Thr	Leu	Ile	Ser	Arg	260	265	270
Ala	Pro	Val	Val	Thr	His	Thr	Tyr	Leu	Glu	Pro	Gly	Pro	Val	Thr	Ala	275	280	285
Gln	Val	Val	Leu	Gln	Ala	Ala	Ile	Pro	Leu	Thr	Ser	Cys	Gly	Ser	Ser	290	295	300
Pro	Val	Pro	Gly	Thr	Thr	Asp	Gly	His	Arg	Pro	Thr	Ala	Glu	Ala	Pro	305	310	315
Asn	Thr	Thr	Ala	Gly	Gln	Val	Pro	Thr	Thr	Glu	Val	Val	Gly	Thr	Thr	325	330	335
Pro	Gly	Gln	Ala	Pro	Thr	Ala	Glu	Pro	Ser	Gly	Thr	Thr	Ser	Val	Gln	340	345	350
Val	Pro	Thr	Thr	Glu	Val	Ile	Ser	Thr	Ala	Pro	Val	Gln	Met	Pro	Thr	355	360	365
Ala	Glu	Ser	Thr	Gly	Met	Thr	Pro	Glu	Lys	Val	Pro	Val	Ser	Glu	Val	370	375	380
Met	Gly	Thr	Thr	Leu	Ala	Glu	Met	Ser	Thr	Pro	Glu	Ala	Thr	Gly	Met	385	390	395
Thr	Pro	Ala	Glu	Val	Ser	Ile	Val	Val	Leu	Ser	Gly	Thr	Thr	Ala	Ala	405	410	415
Gln	Val	Thr	Thr	Thr	Glu	Trp	Val	Glu	Thr	Thr	Ala	Arg	Glu	Leu	Pro	420	425	430
Ile	Pro	Glu	Pro	Glu	Gly	Pro	Asp	Ala	Ser	Ser	Ile	Met	Ser	Thr	Glu	435	440	445
Ser	Ile	Thr	Gly	Ser	Leu	Gly	Pro	Leu	Leu	Asp	Gly	Thr	Ala	Thr	Leu	450	455	460
Arg	Leu	Val	Lys	Arg	Gln	Val	Pro	Leu	Asp	Cys	Val	Leu	Tyr	Arg	Tyr	465	470	475
Gly	Ser	Phe	Ser	Val	Thr	Leu	Asp	Ile	Val	Gln	Gly	Ile	Glu	Ser	Ala	485	490	495
Glu	Ile	Leu	Gln	Ala	Val	Pro	Ser	Gly	Glu	Gly	Asp	Ala	Phe	Glu	Leu	500	505	510
Thr	Val	Ser	Cys	Gln	Gly	Gly	Leu	Pro	Lys	Glu	Ala	Cys	Met	Glu	Ile	515	520	525
Ser	Ser	Pro	Gly	Cys	Gln	Pro	Pro	Ala	Gln	Arg	Leu	Cys	Gln	Pro	Val	530	535	540
Leu	Pro	Ser	Pro	Ala	Cys	Gln	Leu	Val	Leu	His	Gln	Ile	Leu	Lys	Gly	545	550	555
Gly	Ser	Gly	Thr	Tyr	Cys	Leu	Asn	Val	Ser	Leu	Ala	Asp	Thr	Asn	Ser	565	570	575
Leu	Ala	Val	Val	Ser	Thr	Gln	Leu	Ile	Met	Pro	Gly	Gln	Glu	Ala	Gly	580	585	590
Leu	Gly	Gln	Val	Pro	Leu	Ile	Val	Gly	Ile	Leu	Leu	Val	Leu	Met	Ala	595	600	605
Val	Val	Leu	Ala	Ser	Leu	Ile	Tyr	Arg	Arg	Arg	Leu	Met	Lys	Gln	Asp	610	615	620
Phe	Ser	Val	Pro	Gln	Leu	Pro	His	Ser	Ser	Ser	His	Trp	Leu	Arg	Leu	625	630	635
Pro	Arg	Ile	Phe	Cys	Ser	Cys	Pro	Ile	Gly	Glu	Asn	Ser	Pro	Leu	Leu	645	650	655
Ser	Gly	Gln	Gln	Val														

<210> 71
 <211> 309
 <212> PRT
 <213> Homo sapiens

<400> 71

Met	Ser	Leu	Glu	Gln	Arg	Ser	Leu	His	Cys	Lys	Pro	Glu	Glu	Ala	Leu
1				5					10					15	
Glu	Ala	Gln	Gln	Glu	Ala	Leu	Gly	Leu	Val	Cys	Val	Gln	Ala	Ala	Thr
		20					25					30			
Ser	Ser	Ser	Ser	Pro	Leu	Val	Leu	Gly	Thr	Leu	Glu	Glu	Val	Pro	Thr
		35					40					45			
Ala	Gly	Ser	Thr	Asp	Pro	Pro	Gln	Ser	Pro	Gln	Gly	Ala	Ser	Ala	Phe
	50					55					60				
Pro	Thr	Thr	Ile	Asn	Phe	Thr	Arg	Gln	Arg	Gln	Pro	Ser	Glu	Gly	Ser
65				70						75					80
Ser	Ser	Arg	Glu	Glu	Glu	Gly	Pro	Ser	Thr	Ser	Cys	Ile	Leu	Glu	Ser
			85						90					95	
Leu	Phe	Arg	Ala	Val	Ile	Thr	Lys	Lys	Val	Ala	Asp	Leu	Val	Gly	Phe
			100					105					110		
Leu	Leu	Leu	Lys	Tyr	Arg	Ala	Arg	Glu	Pro	Val	Thr	Lys	Ala	Glu	Met
		115					120					125			
Leu	Glu	Ser	Val	Ile	Lys	Asn	Tyr	Lys	His	Cys	Phe	Pro	Glu	Ile	Phe
	130				135						140				
Gly	Lys	Ala	Ser	Glu	Ser	Leu	Gln	Leu	Val	Phe	Gly	Ile	Asp	Val	Lys
145				150						155					160
Glu	Ala	Asp	Pro	Thr	Gly	His	Ser	Tyr	Val	Leu	Val	Thr	Cys	Leu	Gly
			165						170					175	
Leu	Ser	Tyr	Asp	Gly	Leu	Leu	Gly	Asp	Asn	Gln	Ile	Met	Pro	Lys	Thr
		180						185					190		
Gly	Phe	Leu	Ile	Ile	Val	Leu	Val	Met	Ile	Ala	Met	Glu	Gly	Gly	His
		195					200						205		
Ala	Pro	Glu	Glu	Glu	Ile	Trp	Glu	Glu	Leu	Ser	Val	Met	Glu	Val	Tyr
		210				215					220				
Asp	Gly	Arg	Glu	His	Ser	Ala	Tyr	Gly	Glu	Pro	Arg	Lys	Leu	Leu	Thr
225				230						235					240
Gln	Asp	Leu	Val	Gln	Glu	Lys	Tyr	Leu	Glu	Tyr	Arg	Gln	Val	Pro	Asp
			245						250					255	
Ser	Asp	Pro	Ala	Arg	Tyr	Glu	Phe	Leu	Trp	Gly	Pro	Arg	Ala	Leu	Ala
		260						265					270		
Glu	Thr	Ser	Tyr	Val	Lys	Val	Leu	Glu	Tyr	Val	Ile	Lys	Val	Ser	Ala
		275					280					285			
Arg	Val	Arg	Phe	Phe	Phe	Pro	Ser	Leu	Arg	Glu	Ala	Ala	Leu	Arg	Glu
	290					295					300				
Glu	Glu	Glu	Gly	Val											
305															

<210> 72
 <211> 314
 <212> PRT
 <213> Homo sapiens

<400> 72

Met	Pro	Leu	Glu	Gln	Arg	Ser	Gln	His	Cys	Lys	Pro	Glu	Glu	Gly	Leu
1				5					10					15	
Glu	Ala	Arg	Gly	Glu	Ala	Leu	Gly	Leu	Val	Gly	Ala	Gln	Ala	Pro	Ala
			20					25					30		
Thr	Glu	Glu	Gln	Gln	Thr	Ala	Ser	Ser	Ser	Ser	Thr	Leu	Val	Glu	Val
			35				40					45			
Thr	Leu	Gly	Glu	Val	Pro	Ala	Ala	Asp	Ser	Pro	Ser	Pro	Pro	His	Ser
			50			55					60				
Pro	Gln	Gly	Ala	Ser	Ser	Phe	Ser	Thr	Thr	Ile	Asn	Tyr	Thr	Leu	Trp
65					70					75					80
Arg	Gln	Ser	Asp	Glu	Gly	Ser	Ser	Asn	Gln	Glu	Glu	Glu	Gly	Pro	Arg
				85				90						95	
Met	Phe	Pro	Asp	Leu	Glu	Ser	Glu	Phe	Gln	Ala	Ala	Ile	Ser	Arg	Lys
			100					105					110		
Met	Val	Glu	Leu	Val	His	Phe	Leu	Leu	Lys	Tyr	Arg	Ala	Arg	Glu	
			115				120						125		
Pro	Val	Thr	Lys	Ala	Glu	Met	Leu	Glu	Ser	Val	Leu	Arg	Asn	Cys	Gln
			130			135					140				
Asp	Phe	Phe	Pro	Val	Ile	Phe	Ser	Lys	Ala	Ser	Glu	Tyr	Leu	Gln	Leu
145					150					155					160
Val	Phe	Gly	Ile	Glu	Val	Val	Glu	Val	Val	Pro	Ile	Ser	His	Leu	Tyr
				165					170					175	
Ile	Leu	Val	Thr	Cys	Leu	Gly	Leu	Ser	Tyr	Asp	Gly	Leu	Leu	Gly	Asp
			180					185					190		
Asn	Gln	Val	Met	Pro	Lys	Thr	Gly	Leu	Leu	Ile	Ile	Val	Leu	Ala	Ile
			195				200						205		
Ile	Ala	Ile	Glu	Gly	Asp	Cys	Ala	Pro	Glu	Glu	Lys	Ile	Trp	Glu	Glu
			210			215					220				
Leu	Ser	Met	Leu	Glu	Val	Phe	Glu	Gly	Arg	Glu	Asp	Ser	Val	Phe	Ala
225					230					235					240
His	Pro	Arg	Lys	Leu	Leu	Met	Gln	Asp	Leu	Val	Gln	Glu	Asn	Tyr	Leu
				245				250						255	
Glu	Tyr	Arg	Gln	Val	Pro	Gly	Ser	Asp	Pro	Ala	Cys	Tyr	Glu	Phe	Leu
			260					265					270		
Trp	Gly	Pro	Arg	Ala	Leu	Ile	Glu	Thr	Ser	Tyr	Val	Lys	Val	Leu	His
			275				280						285		
His	Thr	Leu	Lys	Ile	Gly	Gly	Glu	Pro	His	Ile	Ser	Tyr	Pro	Pro	Leu
			290			295					300				
His	Glu	Arg	Ala	Leu	Arg	Glu	Gly	Glu	Glu						
305					310										

<210> 73

<211> 314

<212> PRT

<213> Homo sapiens

<400> 73

Met	Pro	Leu	Glu	Gln	Arg	Ser	Gln	His	Cys	Lys	Pro	Glu	Glu	Gly	Leu
1				5					10					15	
Glu	Ala	Arg	Gly	Glu	Ala	Leu	Gly	Leu	Val	Gly	Ala	Gln	Ala	Pro	Ala
			20					25					30		
Thr	Glu	Glu	Gln	Glu	Ala	Ala	Ser	Ser	Ser	Ser	Thr	Leu	Val	Glu	Val
			35				40					45			
Thr	Leu	Gly	Glu	Val	Pro	Ala	Ala	Glu	Ser	Pro	Asp	Pro	Pro	Gln	Ser
			50			55					60				
Pro	Gln	Gly	Ala	Ser	Ser	Leu	Pro	Thr	Thr	Met	Asn	Tyr	Pro	Leu	Trp

65					70					75				80
Ser	Gln	Ser	Tyr	Glu	Asp	Ser	Ser	Asn	Gln	Glu	Glu	Glu	Gly	Pro
				85					90					95
Thr	Phe	Pro	Asp	Leu	Glu	Ser	Glu	Phe	Gln	Ala	Ala	Leu	Ser	Arg
			100					105					110	Lys
Val	Ala	Glu	Leu	Val	His	Phe	Leu	Leu	Leu	Lys	Tyr	Arg	Ala	Arg
		115					120					125		Glu
Pro	Val	Thr	Lys	Ala	Glu	Met	Leu	Gly	Ser	Val	Val	Gly	Asn	Trp
		130				135					140			Gln
Tyr	Phe	Phe	Pro	Val	Ile	Phe	Ser	Lys	Ala	Ser	Ser	Ser	Leu	Gln
145					150				155					160
Val	Phe	Gly	Ile	Glu	Leu	Met	Glu	Val	Asp	Pro	Ile	Gly	His	Leu
			165						170					Tyr
Ile	Phe	Ala	Thr	Cys	Leu	Gly	Leu	Ser	Tyr	Asp	Gly	Leu	Leu	Gly
			180					185					190	Asp
Asn	Gln	Ile	Met	Pro	Lys	Ala	Gly	Leu	Leu	Ile	Ile	Val	Leu	Ala
		195					200					205		Ile
Ile	Ala	Arg	Glu	Gly	Asp	Cys	Ala	Pro	Glu	Glu	Lys	Ile	Trp	Glu
	210				215						220			Glu
Leu	Ser	Val	Leu	Glu	Val	Phe	Glu	Gly	Arg	Glu	Asp	Ser	Ile	Leu
225					230					235				Gly
Asp	Pro	Lys	Lys	Leu	Leu	Thr	Gln	His	Phe	Val	Gln	Glu	Asn	Tyr
			245						250				255	Leu
Glu	Tyr	Arg	Gln	Val	Pro	Gly	Ser	Asp	Pro	Ala	Cys	Tyr	Glu	Phe
		260					265						270	Leu
Trp	Gly	Pro	Arg	Ala	Leu	Val	Glu	Thr	Ser	Tyr	Val	Lys	Val	Leu
	275					280						285		His
His	Met	Val	Lys	Ile	Ser	Gly	Gly	Pro	His	Ile	Ser	Tyr	Pro	Pro
	290				295						300			Leu
His	Glu	Trp	Val	Leu	Arg	Glu	Gly	Glu	Glu					
305					310									

<210> 74
 <211> 180
 <212> PRT
 <213> Homo sapiens

<400> 74
 Met Gln Ala Glu Gly Arg Gly Thr Gly Gly Ser Thr Gly Asp Ala Asp
 1 5 10 15
 Gly Pro Gly Gly Pro Gly Ile Pro Asp Gly Pro Gly Gly Asn Ala Gly
 20 25 30
 Gly Pro Gly Glu Ala Gly Ala Thr Gly Gly Arg Gly Pro Arg Gly Ala
 35 40 45
 Gly Ala Ala Arg Ala Ser Gly Pro Gly Gly Gly Ala Pro Arg Gly Pro
 50 55 60
 His Gly Gly Ala Ala Ser Gly Leu Asn Gly Cys Cys Arg Cys Gly Ala
 65 70 75 80
 Arg Gly Pro Glu Ser Arg Leu Leu Glu Phe Tyr Leu Ala Met Pro Phe
 85 90 95
 Ala Thr Pro Met Glu Ala Glu Leu Ala Arg Arg Ser Leu Ala Gln Asp
 100 105 110
 Ala Pro Pro Leu Pro Val Pro Gly Val Leu Leu Lys Glu Phe Thr Val
 115 120 125
 Ser Gly Asn Ile Leu Thr Ile Arg Leu Thr Ala Ala Asp His Arg Gln
 130 135 140

Leu Gln Leu Ser Ile Ser Ser Cys Leu Gln Gln Leu Ser Leu Leu Met
 145 150 155 160
 Trp Ile Thr Gln Cys Phe Leu Pro Val Phe Leu Ala Gln Pro Pro Ser
 165 170 175
 Gly Gln Arg Arg
 180

<210> 75
 <211> 180
 <212> PRT
 <213> Homo sapiens

<400> 75
 Met Gln Ala Glu Gly Arg Gly Thr Gly Gly Ser Thr Gly Asp Ala Asp
 1 5 10 15
 Gly Pro Gly Gly Pro Gly Ile Pro Asp Gly Pro Gly Gly Asn Ala Gly
 20 25 30
 Gly Pro Gly Glu Ala Gly Ala Thr Gly Gly Arg Gly Pro Arg Gly Ala
 35 40 45
 Gly Ala Ala Arg Ala Ser Gly Pro Arg Gly Gly Ala Pro Arg Gly Pro
 50 55 60
 His Gly Gly Ala Ala Ser Ala Gln Asp Gly Arg Cys Pro Cys Gly Ala
 65 70 75 80
 Arg Arg Pro Asp Ser Arg Leu Leu Glu Leu His Ile Thr Met Pro Phe
 85 90 95
 Ser Ser Pro Met Glu Ala Glu Leu Val Arg Arg Ile Leu Ser Arg Asp
 100 105 110
 Ala Ala Pro Leu Pro Arg Pro Gly Ala Val Leu Lys Asp Phe Thr Val
 115 120 125
 Ser Gly Asn Leu Leu Phe Ile Arg Leu Thr Ala Ala Asp His Arg Gln
 130 135 140
 Leu Gln Leu Ser Ile Ser Ser Cys Leu Gln Gln Leu Ser Leu Leu Met
 145 150 155 160
 Trp Ile Thr Gln Cys Phe Leu Pro Val Phe Leu Ala Gln Ala Pro Ser
 165 170 175
 Gly Gln Arg Arg
 180

<210> 76
 <211> 210
 <212> PRT
 <213> Homo sapiens

<400> 76
 Met Gln Ala Glu Gly Arg Gly Thr Gly Gly Ser Thr Gly Asp Ala Asp
 1 5 10 15
 Gly Pro Gly Gly Pro Gly Ile Pro Asp Gly Pro Gly Gly Asn Ala Gly
 20 25 30
 Gly Pro Gly Glu Ala Gly Ala Thr Gly Gly Arg Gly Pro Arg Gly Ala
 35 40 45
 Gly Ala Ala Arg Ala Ser Gly Pro Arg Gly Gly Ala Pro Arg Gly Pro
 50 55 60
 His Gly Gly Ala Ala Ser Ala Gln Asp Gly Arg Cys Pro Cys Gly Ala
 65 70 75 80
 Arg Arg Pro Asp Ser Arg Leu Leu Glu Leu His Ile Thr Met Pro Phe

Gly	Gln	Met	Ile	Asn	Leu	Arg	Arg	Leu	Leu	Leu	Ser	His	Ile	His	Ala	
			260					265					270			
Ser	Ser	Tyr	Ile	Ser	Pro	Glu	Lys	Glu	Glu	Gln	Tyr	Ile	Ala	Gln	Phe	
		275					280					285				
Thr	Ser	Gln	Phe	Leu	Ser	Leu	Gln	Cys	Leu	Gln	Ala	Leu	Tyr	Val	Asp	
		290				295					300					
Ser	Leu	Phe	Phe	Leu	Arg	Gly	Arg	Leu	Asp	Gln	Leu	Leu	Arg	His	Val	
305					310					315					320	
Met	Asn	Pro	Leu	Glu	Thr	Leu	Ser	Ile	Thr	Asn	Cys	Arg	Leu	Ser	Glu	
				325					330						335	
Gly	Asp	Val	Met	His	Leu	Ser	Gln	Ser	Pro	Ser	Val	Ser	Gln	Leu	Ser	
		340						345					350			
Val	Leu	Ser	Leu	Ser	Gly	Val	Met	Leu	Thr	Asp	Val	Ser	Pro	Glu	Pro	
		355					360					365				
Leu	Gln	Ala	Leu	Leu	Glu	Arg	Ala	Ser	Ala	Thr	Leu	Gln	Asp	Leu	Val	
		370				375					380					
Phe	Asp	Glu	Cys	Gly	Ile	Thr	Asp	Asp	Gln	Leu	Leu	Ala	Leu	Leu	Pro	
385					390				395						400	
Ser	Leu	Ser	His	Cys	Ser	Gln	Leu	Thr	Thr	Leu	Ser	Phe	Tyr	Gly	Asn	
			405						410					415		
Ser	Ile	Ser	Ile	Ser	Ala	Leu	Gln	Ser	Leu	Leu	Gln	His	Leu	Ile	Gly	
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Glu	Asp	Ile	His	Gly	Thr	Leu	His	Leu	Glu	Arg	Leu	Ala	Tyr	Leu	His	
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Ala	Arg	Leu	Arg	Glu	Leu	Leu	Cys	Glu	Leu	Gly	Arg	Pro	Ser	Met	Val	
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Trp	Leu	Ser	Ala	Asn	Pro	Cys	Pro	His	Cys	Gly	Asp	Arg	Thr	Phe	Tyr	
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<400> 78

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Lys	His	Ser	Gln	Pro	Trp	Gln	Val	Leu	Val	Ala	Ser	Arg	Gly	Arg	Ala	
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Phe	His	Pro	Glu	Asp	Thr	Gly	Gln	Val	Phe	Gln	Val	Ser	His	Ser	Phe	
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Gly Ile Thr Ser Trp Gly Ser Glu Pro Cys Ala Leu Pro Glu Arg Pro				
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 <213> Homo sapiens

<400> 79
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50 55 60
Gly Cys Ser Leu Asn Cys Val Asp Asp Ser Gln Asp Tyr Tyr Val Gly
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<210> 84
<211> 752
<212> DNA
<213> Homo sapiens

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<210> 85
<211> 2148
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
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<223> n = A,T,C or G

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 <211> 1466
 <212> DNA
 <213> Homo sapiens

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<210> 87
<211> 990
<212> DNA
<213> Homo sapiens

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<220>
<221> misc_feature
<222> (1)...(990)
<223> n = A,T,C or G

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<210> 88
<211> 702
<212> PRT
<213> Homo sapiens

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<400> 88
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35 40 45
Lys Glu Val Leu Leu Leu Val His Asn Leu Pro Gln His Leu Phe Gly
50 55 60
Tyr Ser Trp Tyr Lys Gly Glu Arg Val Asp Gly Asn Arg Gln Ile Ile
65 70 75 80
Gly Tyr Val Ile Gly Thr Gln Gln Ala Thr Pro Gly Pro Ala Tyr Ser
85 90 95
Gly Arg Glu Ile Ile Tyr Pro Asn Ala Ser Leu Leu Ile Gln Asn Ile
100 105 110
Ile Gln Asn Asp Thr Gly Phe Tyr Thr Leu His Val Ile Lys Ser Asp

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Pro	Asp	Thr	Pro	Ile	Ile	Ser	Pro	Pro	Asp	Ser	Ser	Tyr	Leu	Ser	Gly
		595					600					605			
Ala	Asn	Leu	Asn	Leu	Ser	Cys	His	Ser	Ala	Ser	Asn	Pro	Ser	Pro	Gln
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Tyr	Ser	Trp	Arg	Ile	Asn	Gly	Ile	Pro	Gln	Gln	His	Thr	Gln	Val	Leu
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Phe	Ile	Ala	Lys	Ile	Thr	Pro	Asn	Asn	Asn	Gly	Thr	Tyr	Ala	Cys	Phe
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			660					665					670		
Thr	Val	Ser	Ala	Ser	Gly	Thr	Ser	Pro	Gly	Leu	Ser	Ala	Gly	Ala	Thr
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<210> 89
 <211> 2974
 <212> DNA
 <213> Homo sapiens

<400> 89

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<210> 90
 <211> 1255
 <212> PRT
 <213> Homo sapiens

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Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His
 35          40          45
Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr
 50          55          60
Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val
 65          70          75          80
Gln Gly Tyr Val Leu Ile Ala His Asn Gln Val Arg Gln Val Pro Leu
 85          90          95
Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Asn Tyr
100          105          110
Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Asn Asn Thr Thr Pro
115          120          125
Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln Leu Arg Ser
130          135          140
Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Gln Arg Asn Pro Gln
145          150          155          160
Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe His Lys Asn
165          170          175
Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser Arg Ala Cys
180          185          190
His Pro Cys Ser Pro Met Cys Lys Gly Ser Arg Cys Trp Gly Glu Ser
195          200          205
Ser Glu Asp Cys Gln Ser Leu Thr Arg Thr Val Cys Ala Gly Gly Cys
210          215          220
Ala Arg Cys Lys Gly Pro Leu Pro Thr Asp Cys Cys His Glu Gln Cys
225          230          235          240
Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu Ala Cys Leu
245          250          255
His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro Ala Leu Val

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Arg Lys Val Lys Val Leu Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys
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 Gly Ile Trp Ile Pro Asp Gly Glu Asn Val Lys Ile Pro Val Ala Ile
 740 745 750
 Lys Val Leu Arg Glu Asn Thr Ser Pro Lys Ala Asn Lys Glu Ile Leu
 755 760 765
 Asp Glu Ala Tyr Val Met Ala Gly Val Gly Ser Pro Tyr Val Ser Arg
 770 775 780
 Leu Leu Gly Ile Cys Leu Thr Ser Thr Val Gln Leu Val Thr Gln Leu
 785 790 795 800
 Met Pro Tyr Gly Cys Leu Leu Asp His Val Arg Glu Asn Arg Gly Arg
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 Leu Gly Ser Gln Asp Leu Leu Asn Trp Cys Met Gln Ile Ala Lys Gly
 820 825 830
 Met Ser Tyr Leu Glu Asp Val Arg Leu Val His Arg Asp Leu Ala Ala
 835 840 845
 Arg Asn Val Leu Val Lys Ser Pro Asn His Val Lys Ile Thr Asp Phe
 850 855 860
 Gly Leu Ala Arg Leu Leu Asp Ile Asp Glu Thr Glu Tyr His Ala Asp
 865 870 875 880
 Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu Glu Ser Ile Leu Arg
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 Arg Arg Phe Thr His Gln Ser Asp Val Trp Ser Tyr Gly Val Thr Val
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 Trp Glu Leu Met Thr Phe Gly Ala Lys Pro Tyr Asp Gly Ile Pro Ala
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 930 935 940
 Pro Ile Cys Thr Ile Asp Val Tyr Met Ile Met Val Lys Cys Trp Met
 945 950 955 960
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 965 970 975
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 980 985 990
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 995 1000 1005
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 1045 1050 1055
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 1060 1065 1070
 Ser Pro Leu Ala Pro Ser Glu Gly Ala Gly Ser Asp Val Phe Asp Gly
 1075 1080 1085
 Asp Leu Gly Met Gly Ala Ala Lys Gly Leu Gln Ser Leu Pro Thr His
 1090 1095 1100
 Asp Pro Ser Pro Leu Gln Arg Tyr Ser Glu Asp Pro Thr Val Pro Leu
 1105 1110 1115 1120
 Pro Ser Glu Thr Asp Gly Tyr Val Ala Pro Leu Thr Cys Ser Pro Gln
 1125 1130 1135
 Pro Glu Tyr Val Asn Gln Pro Asp Val Arg Pro Gln Pro Pro Ser Pro
 1140 1145 1150
 Arg Glu Gly Pro Leu Pro Ala Ala Arg Pro Ala Gly Ala Thr Leu Glu
 1155 1160 1165
 Arg Ala Lys Thr Leu Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val

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Phe Ala Phe Gly Gly Ala Val Glu Asn Pro Glu Tyr Leu Thr Pro Gln		
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Gly Gly Ala Ala Pro Gln Pro His Pro Pro Pro Ala Phe Ser Pro Ala		1200
1205	1210	1215
Phe Asp Asn Leu Tyr Tyr Trp Asp Gln Asp Pro Pro Glu Arg Gly Ala		
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Pro Pro Ser Thr Phe Lys Gly Thr Pro Thr Ala Glu Asn Pro Glu Tyr		
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Leu Gly Leu Asp Val Pro Val		
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<210> 91
 <211> 4530
 <212> DNA
 <213> Homo sapiens

<400> 91

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<210> 92
 <211> 976
 <212> PRT
 <213> Homo sapiens

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<400> 92
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Thr Phe Phe Lys Ser Phe Asn Lys Cys Thr Glu Asp Asp Leu Glu Phe
          35             40             45
Pro Phe Ala Lys Thr Asn Leu Ser Lys Asn Gly Glu Asn Ile Asp Ser
          50             55             60
Asp Pro Ala Leu Gln Lys Val Asn Phe Leu Pro Val Leu Glu Gln Val
          65             70             75             80
Gly Asn Ser Asp Cys His Tyr Gln Glu Gly Leu Lys Asp Ser Asp Leu
          85             90             95

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Glu	Asn	Ser	Glu	Gly	Leu	Ser	Arg	Val	Phe	Ser	Lys	Leu	Tyr	Lys	Glu	100	105	110
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Gln	Lys	Glu	Ser	Lys	Leu	Gln	Glu	Asn	Arg	Lys	Ile	Ile	Glu	Ala	Gln	130	135	140
Arg	Lys	Ala	Ile	Gln	Glu	Leu	Gln	Phe	Gly	Asn	Glu	Lys	Val	Ser	Leu	145	150	155
Lys	Leu	Glu	Glu	Gly	Ile	Gln	Glu	Asn	Lys	Asp	Leu	Ile	Lys	Glu	Asn	165	170	175
Asn	Ala	Thr	Arg	His	Leu	Cys	Asn	Leu	Leu	Lys	Glu	Thr	Cys	Ala	Arg	180	185	190
Ser	Ala	Glu	Lys	Thr	Lys	Lys	Tyr	Glu	Tyr	Glu	Arg	Glu	Glu	Thr	Arg	195	200	205
Gln	Val	Tyr	Met	Asp	Leu	Asn	Asn	Asn	Ile	Glu	Lys	Met	Ile	Thr	Ala	210	215	220
His	Gly	Glu	Leu	Arg	Val	Gln	Ala	Glu	Asn	Ser	Arg	Leu	Glu	Met	His	225	230	235
Phe	Lys	Leu	Lys	Glu	Asp	Tyr	Glu	Lys	Ile	Gln	His	Leu	Glu	Gln	Glu	245	250	255
Tyr	Lys	Lys	Glu	Ile	Asn	Asp	Lys	Glu	Lys	Gln	Val	Ser	Leu	Leu	Leu	260	265	270
Ile	Gln	Ile	Thr	Glu	Lys	Glu	Asn	Lys	Met	Lys	Asp	Leu	Thr	Phe	Leu	275	280	285
Leu	Glu	Glu	Ser	Arg	Asp	Lys	Val	Asn	Gln	Leu	Glu	Glu	Lys	Thr	Lys	290	295	300
Leu	Gln	Ser	Glu	Asn	Leu	Lys	Gln	Ser	Ile	Glu	Lys	Gln	His	His	Leu	305	310	315
Thr	Lys	Glu	Leu	Glu	Asp	Ile	Lys	Val	Ser	Leu	Gln	Arg	Ser	Val	Ser	325	330	335
Thr	Gln	Lys	Ala	Leu	Glu	Glu	Asp	Leu	Gln	Ile	Ala	Thr	Lys	Thr	Ile	340	345	350
Cys	Gln	Leu	Thr	Glu	Glu	Lys	Glu	Thr	Gln	Met	Glu	Glu	Ser	Asn	Lys	355	360	365
Ala	Arg	Ala	Ala	His	Ser	Phe	Val	Val	Thr	Glu	Phe	Glu	Thr	Thr	Val	370	375	380
Cys	Ser	Leu	Glu	Glu	Leu	Leu	Arg	Thr	Glu	Gln	Arg	Leu	Glu	Lys		385	390	395
Asn	Glu	Asp	Gln	Leu	Lys	Ile	Leu	Thr	Met	Glu	Leu	Gln	Lys	Lys	Ser	405	410	415
Ser	Glu	Leu	Glu	Glu	Met	Thr	Lys	Leu	Thr	Asn	Asn	Lys	Glu	Val	Glu	420	425	430
Leu	Glu	Glu	Leu	Lys	Lys	Val	Leu	Gly	Glu	Lys	Glu	Thr	Leu	Leu	Tyr	435	440	445
Glu	Asn	Lys	Gln	Phe	Glu	Lys	Ile	Ala	Glu	Glu	Leu	Lys	Gly	Thr	Glu	450	455	460
Gln	Glu	Leu	Ile	Gly	Leu	Leu	Gln	Ala	Arg	Glu	Lys	Glu	Val	His	Asp	465	470	475
Leu	Glu	Ile	Gln	Leu	Thr	Ala	Ile	Thr	Thr	Ser	Glu	Gln	Tyr	Tyr	Ser	485	490	495
Lys	Glu	Val	Lys	Asp	Leu	Lys	Thr	Glu	Leu	Glu	Asn	Glu	Lys	Leu	Lys	500	505	510
Asn	Thr	Glu	Leu	Thr	Ser	His	Cys	Asn	Lys	Leu	Ser	Leu	Glu	Asn	Lys	515	520	525
Glu	Leu	Thr	Gln	Glu	Thr	Ser	Asp	Met	Thr	Leu	Glu	Leu	Lys	Asn	Gln	530	535	540
Gln	Glu	Asp	Ile	Asn	Asn	Asn	Lys	Lys	Gln	Glu	Glu	Arg	Met	Leu	Lys			

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Glu	Tyr	Val	Arg	Glu	Glu	Leu	Lys	Gln	Lys	Arg	Asp	Glu	Val	Lys	Cys
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Lys	Leu	Asp	Lys	Ser	Glu	Glu	Asn	Cys	Asn	Asn	Leu	Arg	Lys	Gln	Val
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Glu	Asn	Lys	Asn	Lys	Tyr	Ile	Glu	Glu	Leu	Gln	Gln	Glu	Asn	Lys	Ala
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Leu	Lys	Lys	Lys	Gly	Thr	Ala	Glu	Ser	Lys	Gln	Leu	Asn	Val	Tyr	Glu
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Ile	Lys	Val	Asn	Lys	Leu	Glu	Leu	Glu	Leu	Glu	Ser	Ala	Lys	Gln	Lys
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Phe	Gly	Glu	Ile	Thr	Asp	Thr	Tyr	Gln	Lys	Glu	Ile	Glu	Asp	Lys	Lys
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Ile	Ser	Glu	Glu	Asn	Leu	Leu	Glu	Glu	Val	Glu	Lys	Ala	Lys	Val	Ile
		675					680					685			
Ala	Asp	Glu	Ala	Val	Lys	Leu	Gln	Lys	Glu	Ile	Asp	Lys	Arg	Cys	Gln
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His	Lys	Ile	Ala	Glu	Met	Val	Ala	Leu	Met	Glu	Lys	His	Lys	His	Gln
705				710						715					720
Tyr	Asp	Lys	Ile	Ile	Glu	Glu	Arg	Asp	Ser	Glu	Leu	Gly	Leu	Tyr	Lys
			725					730					735		
Ser	Lys	Glu	Gln	Glu	Gln	Ser	Ser	Leu	Arg	Ala	Ser	Leu	Glu	Ile	Glu
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Ser	Gln	Thr	Val	Ser	Arg	Asn	Phe	Thr	Ser	Val	Asp	His	Gly	Ile	Ser
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Lys	Asp	Lys	Arg	Asp	Tyr	Leu	Trp	Thr	Ser	Ala	Lys	Asn	Thr	Leu	Ser
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Thr	Pro	Leu	Pro	Lys	Ala	Tyr	Thr	Val	Lys	Thr	Pro	Thr	Lys	Pro	Lys
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Phe	Gly	Ala	Ile	Arg	Lys	Met	Arg	Glu	Asp	Arg	Trp	Ala	Val	Ile	Ala
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<210> 93

<211> 3393

<212> DNA
 <213> Homo sapiens

<400> 93
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 gttcgtacca ccgagatcaa gcagcagtc ggtgtctgcg gtgaaacctc agacctggg 180
 aggcgattcc actttcttca agagtttcaa caaatgtact gaagatgatt tggagtttcc 240
 atttgcaaag actaatctct ccaaaaatgg ggaacacatt gattcagatc ctgctttaca 300
 aaaagttaat ttcttgcccc tgcttgagca ggttggtaat tctgactgtc actatcagga 360
 aggactaaaa gactctgatt tggagaattc agagggattg agcagagtgt tttcaaaact 420
 gtataaggag gctgaaaaga taaaaaatg gaaagtaagt acagaagctg aactgagaca 480
 gaaagaaagt aagttgcaag aaaacagaaa gataattgaa gcacagcgaa aagccattca 540
 ggaactgcaa tttggaaatg aaaaagtaag tttgaaatta gaagaaggaa tacaagaaaa 600
 taaagattta ataaaagaga ataatgccac aaggcattta tgtaatctac tcaaagaaac 660
 ctgtgctaga tctgcagaaa agacaaagaa atatgaatat gaacgggaag aaaccaggca 720
 agtttatatg gatctaaata ataacattga gaaaatgata acagctcatg gggaacttcg 780
 tgtgcaagct gagaattcca gactggaaat gcatttttaag ttaaagggaag attatgaaaa 840
 aatccaacac cttgaacaag aatacaagaa ggaaataaat gacaaggaaa agcaggatc 900
 actactattg atccaaatca ctgagaaaga aaataaaatg aaagatttaa catttctgct 960
 agaggaatcc agagataaag ttaatcaatt agaggaaaag acaaaattac agagtgaaaa 1020
 cttaaaacaa tcaattgaga aacagcatca tttgactaaa gaactagaag atattaaagt 1080
 gtcattacaa agaagtgtga gtactcaaaa ggcttttagag gaagatttac agatagcaac 1140
 aaaaacaatt tgtcagctaa ctgaagaaaa agaaactcaa atggaagaat ctaataaagc 1200
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 catggagctt caaaagaaat caagtgaagc ggaagagatg actaagctta caaataacaa 1380
 agaagtagaa cttgaagaat tgaaaaaagt cttgggagaa aaggaaacac ttttatatga 1440
 aaataaaciaa tttgagaaga ttgctgaaga attaaaagga acagaacaag aactaatttg 1500
 tcttctccaa gccagagaga aagaagtaca tgatttggaa atacagttaa ctgccattac 1560
 cacaagtga cagtattatt caaaagaggt taaagatcta aaaactgagc ttgaaaacga 1620
 gaagcttaag aatactgaat taacttcaca ctgcaacaag ctttcactag aaaacaaaga 1680
 gctcacacag gaaacaagt atatgacctt agaactcaag aatcagcaag aagatattaa 1740
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 aaccaatta agaatgaac tagaatatgt gagagaagag ctaaaacaga aaagagatga 1860
 agttaaatgt aaattggaca agagtgaaga aaattgtaac aatttaagga aacaagttga 1920
 aaataaaaaa aagtatatgt aagaacttca gcaggagaat aaggccttga aaaaaaagg 1980
 tacagcagaa agcaagcaac tgaatgttta tgagataaag gtcaataaat tagagttaga 2040
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 ggacaaaaag atatcagaag aaaatctttt ggaagaggtt gagaaagcaa aagtaatagc 2160
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 aatggtagca cttatggaaa aacataagca ccaatatgat aagatcattg aagaaagaga 2280
 ctcagaatta ggactttata agagcaaaga acaagaacag tcatcactga gagcatcttt 2340
 ggagattgaa ctatccaatc tcaaagctga acttttgtct gttagaagc aacttgaaat 2400
 agaaagagaa gagaaggaaa aactcaaaag agaggcaaaa gaaaacacag ctactcttaa 2460
 agaaaaaaa gacaagaaaa cacaaacatt tttattggaa acacctgaaa tttattggaa 2520
 attggattct aaagcagttc cttcacaaac tgtatctcga aatttcacat cagttgatca 2580
 tggcatatcc aaagataaaa gagactatct gtggacatct gccaaaaata ctttatctac 2640
 accattgcca aaggcatata cagtgaagac accaacaaaa ccaaaactac agcaaagaga 2700
 aaacttgaat ataccattg aagaaagtaa aaaaaagaga aaaatggcct ttgaatttga 2760
 tattaattca gatagttcag aaactactga tcttttgagc atggtttcag aagaagagac 2820
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 aaaaaaggcc cttcatctc taacaacccc tggacctaca ctgaagtttg gagctataag 2940
 aaaaatgagg gaggaccgtt gggctgtaat tgctaaaatg gatagaaaaa aaaaactaaa 3000
 agaagctgaa aagttatttg ttttaattca gagaatcagt gtagttaagg agcctaataa 3060
 cgtgaaactt atagtttaata ttttgttctt atttgccaga gccacatttt atctggaagt 3120
 tgagacttaa aaaatacttg catgaatgat ttgtgtttct ttatattttt agcctaaatg 3180

ttaactacat attgtctgga aacctgtcat tgtattcaga taattagatg attatatatt 3240
 gttgttactt tttcttgat tcatgaaaac tgtttttact aagttttcaa atttgtaaag 3300
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 atattttggg tgcaaaaaaa aaaaaaaaaa aaa 3393

<210> 94
 <211> 188
 <212> PRT
 <213> Homo sapiens

<400> 94
 Met Asn Gly Asp Asp Ala Phe Ala Arg Arg Pro Arg Asp Asp Ala Gln
 1 5 10 15
 Ile Ser Glu Lys Leu Arg Lys Ala Phe Asp Asp Ile Ala Lys Tyr Phe
 20 25 30
 Ser Lys Lys Glu Trp Glu Lys Met Lys Ser Ser Glu Lys Ile Val Tyr
 35 40 45
 Val Tyr Met Lys Leu Asn Tyr Glu Val Met Thr Lys Leu Gly Phe Lys
 50 55 60
 Val Thr Leu Pro Pro Phe Met Arg Ser Lys Arg Ala Ala Asp Phe His
 65 70 75 80
 Gly Asn Asp Phe Gly Asn Asp Arg Asn His Arg Asn Gln Val Glu Arg
 85 90 95
 Pro Gln Met Thr Phe Gly Ser Leu Gln Arg Ile Phe Pro Lys Ile Met
 100 105 110
 Pro Lys Lys Pro Ala Glu Glu Glu Asn Gly Leu Lys Glu Val Pro Glu
 115 120 125
 Ala Ser Gly Pro Gln Asn Asp Gly Lys Gln Leu Cys Pro Pro Gly Asn
 130 135 140
 Pro Ser Thr Leu Glu Lys Ile Asn Lys Thr Ser Gly Pro Lys Arg Gly
 145 150 155 160
 Lys His Ala Trp Thr His Arg Leu Arg Glu Arg Lys Gln Leu Val Val
 165 170 175
 Tyr Glu Glu Ile Ser Asp Pro Glu Glu Asp Asp Glu
 180 185

<210> 95
 <211> 576
 <212> DNA
 <213> Homo sapiens

<400> 95
 atgaacggag acgacgcctt tgcaaggaga cccagggatg atgctcaa atcagagaag 60
 ttacgaaagg cttcgatga tattgccaaa tacttctcta agaaagagtg ggaaaagatg 120
 aaatcctcgg agaaaatcgt ctatgtgtat atgaagctaa actatgaggt catgactaaa 180
 ctaggtttca aggtcaccct cccaccttct atgcgtagta aacgggctgc agacttccac 240
 gggaatgatt ttggtaacga tcgaaaccac aggaatcagg ttgaacgtcc tcagatgact 300
 ttcggcagcc tccagagaat cttcccgaag atcatgccca agaagccagc agaggaagaa 360
 aatggtttga aggaagtgcc agaggcatct ggcccacaaa atgatgggaa acagctgtgc 420
 cccccgggaa atccaagtac cttggagaag attaacaaga catctggacc caaaaggggg 480
 aaacatgcct ggaccacag actgcgtgag agaaagcagc tgggtggtta tgaagagatc 540
 agcgaccctg aggaagatga cgagtaactc ccctcg 576

<210> 96
 <211> 94
 <212> PRT

<213> Homo sapiens

<400> 96

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Pro Ala Thr Gln Arg Gln Asp Pro Ala Ala Ala Gln Glu Gly Glu Asp
 1           5           10           15
Glu Gly Ala Ser Ala Gly Gln Gly Pro Lys Pro Glu Ala Asp Ser Gln
          20           25           30
Glu Gln Gly His Pro Gln Thr Gly Cys Glu Cys Glu Asp Gly Pro Asp
          35           40           45
Gly Gln Glu Met Asp Pro Pro Asn Pro Glu Glu Val Lys Thr Pro Glu
          50           55           60
Glu Glu Met Arg Ser His Tyr Val Ala Gln Thr Gly Ile Leu Trp Leu
65           70           75           80
Leu Met Asn Asn Cys Phe Leu Asn Leu Ser Pro Arg Lys Pro
          85           90
```

<210> 97

<211> 646

<212> DNA

<213> Homo sapiens

<400> 97

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ctgccgtccg gactcttttt cctctactga gattcatctg tgtgaaatat gagttggcga 60
ggaagatcga cctatcggcc tagaccaaga cgctacgtag agcctcctga aatgattggg 120
cctatgcggc ccgagcagtt cagtgatgaa gtggaaccag caacacctga agaaggggaa 180
ccagcaactc aacgtcagga tcctgcagct gctcaggagg gagaggatga gggagcatct 240
gcaggtcaag ggccgaagcc tgaagctgat agccaggaac agggtcaccc acagactggg 300
tgtgagtgtg aagatgggtc tgatgggcag gagatggacc cgccaaatcc agaggagggtg 360
aaaacgcctg aagaagagat gaggtctcac tatgttgccc agactgggat tctctggctt 420
ttaatgaaca attgcttctt aaatctttcc ccacggaaac cttgagtgcac tgaaatatca 480
aatggcgaga gaccgttttag ttcctatcat ctgtggcatg tgaagggcaa tcacagtgtt 540
aaaagaagac atgctgaaat gttgcaggct gctcctatgt tggaaaattc ttcattgaag 600
ttctcccaat aaagctttac agccttctgc aaagaaaaaa aaaaaa 646
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<210> 98

<211> 98

<212> PRT

<213> Homo sapiens

<400> 98

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His Cys Pro Thr Glu Asn Glu Pro Asp Leu Ala Gln Cys Phe Phe Cys
 1           5           10           15
Phe Lys Glu Leu Glu Gly Trp Glu Pro Asp Asp Asp Pro Ile Glu Glu
          20           25           30
His Lys Lys His Ser Ser Gly Cys Ala Phe Leu Ser Val Lys Lys Gln
          35           40           45
Phe Glu Glu Leu Thr Leu Gly Glu Phe Leu Lys Leu Asp Arg Glu Arg
          50           55           60
Ala Lys Asn Lys Ile Ala Lys Glu Thr Asn Asn Lys Lys Lys Glu Phe
65           70           75           80
Glu Glu Thr Ala Lys Lys Val Arg Arg Ala Ile Glu Gln Leu Ala Ala
          85           90           95
Met Asp
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<210> 99
 <211> 1619
 <212> DNA
 <213> Homo sapiens

<400> 99
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 gacgttgccc cctgcctggc agccctttct caaggaccac cgcatctcta cattcaagaa 120
 ctggcccttc ttggagggct gcgcctgcac cccggagcgg atggccgagg ctggcttcat 180
 ccactgcccc actgagaacg agccagactt ggcccagtgt ttcttctgct tcaaggagct 240
 ggaaggctgg gagccagatg acgaccccat agaggaacat aaaaagcatt cgtccggttg 300
 cgctttcctt tctgtcaaga agcagtttga agaattaacc cttggtgaat ttttgaaact 360
 ggacagagaa agagccaaga acaaaattgc aaaggaaacc aacaataaga agaaagaatt 420
 tgaggaaact gcgaagaaag tgccgcgtgc catcgagcag ctggctgcca tggattgagg 480
 cctctggccg gagctgcctg gtcccagagt ggctgcacca cttccagggt ttattccctg 540
 gtgccaccag ccttcctgtg ggccccttag caatgtctta ggaaaggaga tcaacatttt 600
 caaattagat gtttcaactg tgctcctgtt ttgtcttgaa agtggcacca gaggtgcttc 660
 tgctgtgca gcgggtgctg ctggtaacag tggtgcttc tctctctctc tctctttttt 720
 gggggctcat ttttgctgtt ttgattcccg ggcttaccag gtgagaagtg agggaggaag 780
 aaggcagtgt cctttttgct agagctgaca gctttgttcg cgtgggcaga gccttcaca 840
 gtgaatgtgt ctggacctca tgttgttgag gctgtcacag tcctgagtgt ggacttggca 900
 ggtgctgtgt gaatctgagc tgcaggttcc ttatctgtca cactgtgcc tcctcagagg 960
 acagtttttt tggtgtgtgt tttttttgtt tttttttttt ggtagatgca tgacttgtgt 1020
 gtgatgagag aatggagaca ggtccctgg ctcctctact gtttaacaac atggctttct 1080
 tattttgttt gaattgttaa ttcacagaat agcacaact acaattaaa ctaagcaca 1140
 agccattcta agtcattggg gaaacggggt gaacttcagg tggatgagga gacagaatag 1200
 agtgatagga agcgtctggc agatactcct ttggccactg ctgtgtgatt agacaggccc 1260
 agtgagccgc ggggcacatg ctggccgctc ctcctcaga aaaaggcagt ggcctaaatc 1320
 ctttttaaat gacttggctc gatgctgtgg gggactggct gggctgctgc aggcctgtgt 1380
 tctgtcagcc caaccttcac atctgtcacg ttctccacac gggggagaga cgcagtcgc 1440
 ccaggtcccc gctttctttg gaggcagcag ctcgcgcagg gctgaagtct ggcgtaagat 1500
 gatggatttg attcgccctc ctccctgtca tagagctgca ggggtggattg ttacagcttc 1560
 gctggaacc tctggaggtc atctcggtg ttctgagaa ataaaaagcc tgtcatttc 1619

<210> 100
 <211> 74
 <212> PRT
 <213> Homo sapiens

<400> 100
 Cys Trp Tyr Cys Arg Arg Arg Asn Gly Tyr Arg Ala Leu Met Asp Lys
 1 5 10 15
 Ser Leu His Val Gly Thr Gln Cys Ala Leu Thr Arg Arg Cys Pro Gln
 20 25 30
 Glu Gly Phe Asp His Arg Asp Ser Lys Val Ser Leu Gln Glu Lys Asn
 35 40 45
 Cys Glu Pro Val Val Pro Asn Ala Pro Pro Ala Tyr Glu Lys Leu Ser
 50 55 60
 Ala Glu Gln Ser Pro Pro Pro Tyr Ser Pro
 65 70

<210> 101
 <211> 1524
 <212> DNA
 <213> Homo sapiens

<400> 101

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agcagacaga ggactctcat taaggaaggt gtccctgtgcc ctgaccctac aagatgccaa 60
gagaagatgc tcacttcac tcttggtacc ccaagaagg gcacggccac tcttacacca 120
cggctgaaga ggccgctggg atcgcatcc tgacagtgat cctgggagtc ttactgctca 180
tcggctgttg gtattgtaga agacgaaatg gatacagagc cttgatggat aaaagtcttc 240
atgttggcac tcaatgtgcc ttaacaagaa gatgccaca agaagggttt gatcatcggg 300
acagcaaagt gtctcttcaa gagaaaaact gtgaacctgt ggttcccaat gctccacctg 360
cttatgagaa actctctgca gaacagtcac caccacctta ttcaccttaa gagccagcga 420
gacacctgag acatgctgaa attatttctc tcacactttt gcttgaattt aatacagaca 480
tctaattgtt tcctttggaa tgggttagga aaaatgcaag ccatctctaa taataagtca 540
gtgttaaaat tttagtaggt ccgctagcag tactaatcat gtgaggaaat gatgagaaat 600
attaaattgg gaaaactcca tcaataaatg ttgcaatgca tgatactatc tgtgccagag 660
gtaatgttag taaatccatg gtgttatttt ctgagagaca gaattcaagt gggatttctg 720
gggcatcca atttctcttt acttgaaatt tggctaataa caaactagtc aggttttctga 780
accttgaccg acatgaactg tacacagaat tgttccagta ctatggagtg ctcacaaagg 840
atactttttac aggttaagac aaagggttga ctggcctatt tatctgatca agaacatgtc 900
agcaatgtct ctttgtgctc taaaattcta ttatactaca ataatatatt gtaaagatcc 960
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cgcatcttg gctcaccata acctccgcct ccaggttca agcaattctc ctgccttagc 1080
ctcctgagta gctgggatta caggcgtgcg ccactatgcc tgactaattt tgtagtttta 1140
gtagagacgg ggtttctcca tgttggtcag gctgggtctc aactcctgac ctcaggtgat 1200
ctgcccgcct cagcctccca aagtgtgga attacaggc tgagccacca cgcctggctg 1260
gatcctatat cttaggttaag acatataacg cagtctaatt acatttact tcaaggctca 1320
atgctattct aactaatgac aagtattttc tactaaacca gaaattggta gaaggattta 1380
aataagtaaa agctactatg tactgcctta gtgctgatgc ctgtgtactg ccttaaattgt 1440
acctatggca atttagctct cttgggttcc caaatccctc tcacaagaat gtgcagaaga 1500
aatcataaag gatcagagat tctg 1524
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<210> 102

<211> 43

<212> PRT

<213> Homo sapiens

<400> 102

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Met Ala Ala Arg Ala Val Phe Leu Ala Leu Ser Ala Gln Leu Leu Gln
 1           5           10          15
Ala Arg Leu Met Lys Glu Glu Ser Pro Val Val Ser Trp Arg Leu Glu
 20          25          30
Pro Glu Asp Gly Thr Ala Leu Cys Phe Ile Phe
 35          40
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<210> 103

<211> 1004

<212> DNA

<213> Homo sapiens

<400> 103

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cgccaattta gggctctcgg tatctcccgc tgagctgctc tgttcccggc ttagaggacc 60
aggagaaggg ggagctggag gctggagcct gtaacaccgt ggctcgtctc actctggatg 120
gtggtggcaa cagagatggc agcgcagctg gagtgttagg agggcggcct gagcggtagg 180
agtggggctg gagcagtaag atggcgcca gagcggtttt tctggcattg tctgccagc 240
tgctccaagc caggctgatg aaggaggagt cccctgtggt gagctggagg ttggagcctg 300
aagacggcac agctctgtgc ttcactttct gaggttgtgg cagccacggt gatggagacg 360
gcagctcaac aggagcaata ggaggagatg gagtttctact gtgtcagcca ggatgggtctc 420
gatctcctga cctcgtgatc cgcccgcctt ggccttccaa agtgccgaga ttacagcgat 480
gtgcattttg taagcacttt ggagccacta tcaaattgctg tgaagagaaa tgtaccacga 540
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tgtatcatta tccttgtgct gcaggagccg gctcctttca ggatttcagt cacatcttcc 600
tgctttgtcc agaacacatt gaccaagctc ctgaaagatg taagtttact acgcatagac 660
ttttaaaactt caaccaatgt atttactgaa aataacaaat gttgtaaatt ccctgagtgt 720
tattctactt gtattaaaag gtaataatac ataatcatta aaatctgagg gatcattgcc 780
agagattgtt ggggagggaa atgttatcaa cggtttcatt gaaattaaat ccaaaaagtt 840
atttcctcag aaaaatcaaa taaagtttgc atgtttttta ttcttaaaaac attttaaaaa 900
ccactgtaga atgatgtaaa tagggactgt gcagtatttc tgacatatac tataaaatta 960
ttaaaaagtc aatcagtatt caacatcttt tacactaaaa agcc 1004

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<210> 104
<211> 9
<212> PRT
<213> Homo sapiens

```

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<400> 104
Trp Val Leu Thr Ala Ala His Cys Ile
1 5

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<210> 105
<211> 263
<212> PRT
<213> Homo sapiens

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<400> 105
Pro Met Trp Phe Leu Val Leu Cys Leu Ala Leu Ser Leu Gly Gly Thr
1 5 10 15
Gly Ala Ala Pro Pro Ile Gln Ser Arg Ile Val Gly Gly Trp Glu Cys
20 25 30
Glu Gln His Ser Gln Pro Trp Gln Ala Ala Leu Tyr His Phe Ser Thr
35 40 45
Phe Gln Cys Gly Gly Ile Leu Val His Arg Gln Trp Val Leu Thr Ala
50 55 60
Ala His Cys Ile Ser Asp Asn Tyr Gln Leu Trp Leu Gly Arg His Asn
65 70 75 80
Leu Phe Asp Asp Glu Asn Thr Ala Gln Phe Val His Val Ser Glu Ser
85 90 95
Phe Pro His Pro Gly Phe Asn Met Ser Leu Leu Glu Asn His Thr Arg
100 105 110
Gln Ala Asp Glu Asp Tyr Ser His Asp Leu Met Leu Leu Arg Leu Thr
115 120 125
Glu Pro Ala Asp Thr Ile Thr Asp Ala Val Lys Val Val Glu Leu Pro
130 135 140
Thr Gln Glu Pro Glu Val Gly Ser Thr Cys Leu Ala Ser Gly Trp Gly
145 150 155 160
Ser Ile Glu Pro Glu Asn Phe Ser Phe Pro Asp Asp Leu Gln Cys Val
165 170 175
Asp Leu Lys Ile Leu Pro Asn Asp Glu Cys Glu Lys Ala His Val Gln
180 185 190
Lys Val Thr Asp Phe Met Leu Cys Val Gly His Leu Glu Gly Gly Lys
195 200 205
Asp Thr Cys Val Gly Asp Ser Gly Gly Pro Leu Met Cys Asp Gly Val
210 215 220
Leu Gln Gly Val Thr Ser Trp Gly Tyr Val Pro Cys Gly Thr Pro Asn
225 230 235 240
Lys Pro Ser Val Ala Val Arg Val Leu Ser Tyr Val Lys Trp Ile Glu
245 250 255

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Asp Thr Ile Ala Glu Asn Ser
260

<210> 106
<211> 270
<212> PRT
<213> Homo sapiens

<400> 106
Pro Met Ile Arg Thr Leu Leu Leu Ser Thr Leu Val Ala Gly Ala Leu
1 5 10 15
Ser Cys Gly Asp Pro Thr Tyr Pro Pro Tyr Val Thr Arg Val Val Gly
20 25 30
Gly Glu Glu Ala Arg Pro Asn Ser Trp Pro Trp Gln Val Ser Leu Gln
35 40 45
Tyr Ser Ser Asn Gly Lys Trp Tyr His Thr Cys Gly Gly Ser Leu Ile
50 55 60
Ala Asn Ser Trp Val Leu Thr Ala Ala His Cys Ile Ser Ser Ser Arg
65 70 75 80
Thr Tyr Arg Val Gly Leu Gly Arg His Asn Leu Tyr Val Ala Glu Ser
85 90 95
Gly Ser Leu Ala Val Ser Val Ser Lys Ile Val Val His Lys Asp Trp
100 105 110
Asn Ser Asn Gln Ile Ser Lys Gly Asn Asp Ile Ala Leu Leu Lys Leu
115 120 125
Ala Asn Pro Val Ser Leu Thr Asp Lys Ile Gln Leu Ala Cys Leu Pro
130 135 140
Pro Ala Gly Thr Ile Leu Pro Asn Asn Tyr Pro Cys Tyr Val Thr Gly
145 150 155 160
Trp Gly Arg Leu Gln Thr Asn Gly Ala Val Pro Asp Val Leu Gln Gln
165 170 175
Gly Arg Leu Leu Val Val Asp Tyr Ala Thr Cys Ser Ser Ser Ala Trp
180 185 190
Trp Gly Ser Ser Val Lys Thr Ser Met Ile Cys Ala Gly Gly Asp Gly
195 200 205
Val Ile Ser Ser Cys Asn Gly Asp Ser Gly Gly Pro Leu Asn Cys Gln
210 215 220
Ala Ser Asp Gly Arg Trp Gln Val His Gly Ile Val Ser Phe Gly Ser
225 230 235 240
Arg Leu Gly Cys Asn Tyr Tyr His Lys Pro Ser Val Phe Thr Arg Val
245 250 255
Ser Asn Tyr Ile Asp Trp Ile Asn Ser Val Ile Ala Asn Asn
260 265 270

<210> 107
<211> 270
<212> PRT
<213> Homo sapiens

<400> 107
Pro Met Ile Arg Thr Leu Leu Leu Ser Thr Leu Val Ala Gly Ala Leu
1 5 10 15
Ser Cys Gly Val Ser Thr Tyr Ala Pro Asp Met Ser Arg Met Leu Gly
20 25 30
Gly Glu Glu Ala Arg Pro Asn Ser Trp Pro Trp Gln Val Ser Leu Gln

Asp Leu Phe Val Trp Met His Tyr Tyr
1 5

<210> 111
<211> 9
<212> PRT
<213> Homo sapiens

<400> 111
Asp Ala Leu Leu Gly Gly Ser Glu Ile
1 5

<210> 112
<211> 10
<212> PRT
<213> Homo sapiens

<400> 112
Gly Ser Glu Ile Trp Arg Asp Ile Asp Phe
1 5 10

<210> 113
<211> 9
<212> PRT
<213> Homo sapiens

<400> 113
Ser Glu Ile Trp Arg Asp Ile Asp Phe
1 5

<210> 114
<211> 9
<212> PRT
<213> Homo sapiens

<400> 114
Glu Ile Trp Arg Asp Ile Asp Phe Ala
1 5

<210> 115
<211> 10
<212> PRT
<213> Homo sapiens

<400> 115
Leu Gln Glu Val Tyr Pro Glu Ala Asn Ala
1 5 10

<210> 116
<211> 10
<212> PRT

<213> Homosapiens

<400> 116

Glu Val Tyr Pro Glu Ala Asn Ala Pro Ile
1 5 10

<210> 117

<211> 9

<212> PRT

<213> Homosapiens

<400> 117

Val Tyr Pro Glu Ala Asn Ala Pro Ile
1 5

<210> 118

<211> 8

<212> PRT

<213> Homosapiens

<400> 118

Tyr Pro Glu Ala Asn Ala Pro Ile
1 5

<210> 119

<211> 10

<212> PRT

<213> Homosapiens

<400> 119

Tyr Pro Glu Ala Asn Ala Pro Ile Gly His
1 5 10

<210> 120

<211> 10

<212> PRT

<213> Homosapiens

<400> 120

Ala Pro Ile Gly His Asn Arg Glu Ser Tyr
1 5 10

<210> 121

<211> 9

<212> PRT

<213> Homosapiens

<400> 121

Pro Ile Gly His Asn Arg Glu Ser Tyr
1 5

<210> 122
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